

09 / 807, 432

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D rug-resistance genotyping in HIV-1 therapy: the VIRAD APT randomised controlled trial - group of 4 »

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J Durant, P Clevenbergh, P Halfon, P Delgiudice, S ... - THE LANCET, 1999 - bms.com.ve

... For each sample sequenced, the resulting assays were base called with **GeneObjects**

(Visible Genetics), aligned, and assembled together with GeneLibrarian ...

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Detection of Infectious HIV in Circulating Monocytes From Patients on Prolonged Highly Active ... - group of 4 »

O Lambotte, Y Taoufik, MG de Goër, C Wallon, C ... - JAIDS Journal of Acquired Immune Deficiency Syndromes, 2000 - jaids.com

... **Geneobjects** and Genelibrarian software were used for base calling and sequence data analysis (Visible Genetics). RESULTS TOP. HIV ...

Cited by 60 - Related Articles - Cached - Web Search - BL Direct

Sequencing of the protease and reverse transcriptase from HIV-1 subtype A and D, using visible ...

JM Lacroix, RM Lloyd, R Batra, A Bouwvitch, D ... - Abstr Intersci Conf Antimicrob Agents Chemother Intersci ..., 1998 - gateway.nlm.nih.gov

... each portion sequenced. The sequences were base called, aligned and assembled together using **GeneObjects** software. The sequences ...

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R. Tamouza A novel HLA-B null allele (B* 4022N)

K Sadki, V Schaeffer, Z Tatari, J Chabod, A ... - Tissue Antigens, 2000 - ingentaconnect.com

... primers. The sequencing products were analysed on the MicroGene Blaster automated sequencer and interpreted by **GeneObjects** software. ...

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Adherence to protease inhibitors, HIV-1 viral load, and development of drug resistance in an ... - group of 4 »

DR Bangsberg, FM Hecht, ED Charlebois, AR Zolopa, ... - AIDS, 2000 - aidsonline.com

... Chromatograms for each reaction were analysed using **GeneObjects** software (Visible Genetics) and compared with the sequence of HIV-1 HXB2

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Persisting long-term benefit of genotype-guided treatment for HIV-infected patients failing HAART. ... - group of 3 »

P Clevenbergh, J Durant, P Halfon, P del Giudice, ... - Antivir Ther, 2000 - intmedpress.com

... For each sample sequenced, the resulting assays were base called using **GeneObjects** software (Visible Genetics), aligned and assembled together using the ...

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Endoglin Expression Is Reduced in Normal Vessels but Still Detectable in Arteriovenous Malformations ... - group of 3 »

A Bourdeau, U Cymerman, ME Paquet, W Meschino, WC ... - 2000 - ajp.amjpathol.org

... for 30 to 40 minutes. The data were analyzed using **GeneObjects** DNA analysis software (Visible Genetics Inc.). The ratio of the peak ...

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High performance DNA sequencing, and the detection of mutations and polymorphisms, on the Clipper ... - group of 4 »

A Bouevitch, D Chan, K Chan, S Darasch, R ... - Electrophoresis, 1999 - doi.wiley.com

Page 1. High performance DNA sequencing, and the detection of mutations and polymorphisms, on the Clipper sequencer The Visible Genetics ...

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Comparison of twice-daily stavudine plus once-or twice-daily didanosine and nevirapine in early ... - group of 5 »

F Garcia, H Knobel, MA Sambeat, J Arrizabalaga, M ... - AIDS, 2000 - aidsonline.com

... Base-calling was performed using **GeneObjects** software (Version 3.0; Visible Genetics Ltd.), and the bases were aligned and assembled using GeneLibrarian ...

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Identification of Hereditary Hemorrhagic Telangiectasia Type 1 in Newborns by Protein Expression and ... - group of 3 »

U CYMERMAN, S VERA, N PECE-BARBARA, A BOURDEAU, RI ... - Pediatric Research, 2000 - pedresearch.org

... (VGI), Toronto, Ontario, Canada] at 1300 V at 54°C for 30–40 min. The data were analyzed using **GeneObjects** DNA analysis software (VGI). ...

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